# Evolutionary K-means clustering (E-means) using Genetic Algorithms 1.0

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## **Chapter 1**

## evolutionary-clustering

An enhanced highly parallel K-means clustering algorithm using evolutionary strategies to perform metaheuristic optimization.

2	evolutionary-clustering

## Chapter 2

## **Data Structure Index**

2.1	Data Structures	
Here	are the data structures with brief descriptions:	
ne	a state action 64	

**Data Structure Index** 

## **Chapter 3**

## File Index

### 3.1 File List

Here is a list of all files with brief descriptions:

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## **Chapter 4**

## **Data Structure Documentation**

### 4.1 pcg\_state\_setseq\_64 Struct Reference

```
#include <pcg_basic.h>
```

#### **Data Fields**

- uint64\_t state
- uint64\_t inc

#### 4.1.1 Detailed Description

Definition at line 40 of file pcg\_basic.h.

#### 4.1.2 Field Documentation

4.1.2.1 uint64\_t inc

Definition at line 42 of file pcg\_basic.h.

4.1.2.2 uint64\_t state

Definition at line 41 of file pcg\_basic.h.

The documentation for this struct was generated from the following file:

• include/pcg\_basic.h



### **Chapter 5**

### **File Documentation**

#### 5.1 include/cluster.h File Reference

```
#include <gsl/gsl_matrix.h>
#include "pcg_basic.h"
```

#### **Functions**

- int lloyd\_random (int trials, gsl\_matrix \*data, int n\_clusters, gsl\_matrix \*\*clusters, pcg32\_random\_t \*rng)
- int lloyd\_defined (int trials, gsl\_matrix \*centroids, gsl\_matrix \*data, int n\_clusters, gsl\_matrix \*\*clusters)
- int calc\_centroids (gsl\_matrix \*centroids, gsl\_matrix \*data, int n\_clusters, gsl\_matrix \*\*clusters)
- int calc\_bounds (gsl\_matrix \*data, gsl\_matrix \*bounds)
- int random\_centroids (gsl\_matrix \*centroids, gsl\_matrix \*bounds, pcg32\_random\_t \*rng)

#### 5.1.1 Function Documentation

```
5.1.1.1 int calc_bounds ( gsl_matrix * data, gsl_matrix * bounds )
```

Calculates the minimum and maximum bounds based on the data.

#### **Parameters**

data	Point to matrix containing the data
bounds	The min/max bounds for each dimensions of the data

#### Returns

The status code, 0 for SUCCESS, 1 for ERROR

Definition at line 334 of file cluster.c.

5.1.1.2 int calc\_centroids (  $gsl_matrix * centroids$ ,  $gsl_matrix * data$ , int  $n_clusters$ ,  $gsl_matrix ** clusters$  )

Calculate the new centroids using the clustering assignment.

#### **Parameters**

	centroids	Pointer to matrix containing centroids to be updated		
	data	Point to matrix containing the data		
n_clusters The number of clusters				
	cluster	Pointer to vector containing cluster assignment		

#### Returns

The status code, 0 for SUCCESS, 1 for ERROR

Definition at line 307 of file cluster.c.

5.1.1.3 int lloyd\_defined ( int trials,  $gsl_matrix * centroids$ ,  $gsl_matrix * data$ , int  $n_clusters$ ,  $gsl_matrix ** clusters$ )

Performs Lloyd's algorithm using the defined centroids.

#### **Parameters**

trials	Number of trials to perform			
centroids Pointer to matrix containing the centroids				
data	Pointer to matrix containing the data			
n_clusters The number of clusters				
clusters	Pointer to array of matrices containing data in clusters			

#### Returns

The status code, 0 for SUCCESS, 1 for ERROR

Definition at line 191 of file cluster.c.

Here is the call graph for this function:



5.1.1.4 int lloyd\_random ( int trials,  $gsl_matrix * data$ , int  $n_clusters$ ,  $gsl_matrix ** clusters$ ,  $pcg32_random_t * rng$  )

Performs Lloyd's algorithm using random initial centroids.

#### **Parameters**

trials	Number of trials to perform
data	Pointer to matrix containing the data
n_clusters	The number of clusters
clusters	Pointer to array of matrices containing data in clusters

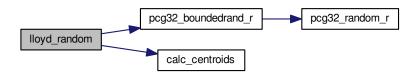
rng	Pointer to the random number generator

#### Returns

The status code, 0 for SUCCESS, 1 for ERROR

Definition at line 33 of file cluster.c.

Here is the call graph for this function:



5.1.1.5 int random\_centroids ( gsl\_matrix \* centroids, gsl\_matrix \* bounds, pcg32\_random\_t \* rng )

Generates random centroids within the bounds for each dimension.

### **Parameters**

centroids	Pointer to matrix containing centroids to be updated
bounds	The min/max bounds for each dimensions of the data
rng	Pointer to the random number generator

#### Returns

The status code, 0 for SUCCESS, 1 for ERROR

Definition at line 362 of file cluster.c.

Here is the call graph for this function:



#### 5.2 include/fitness.h File Reference

#include <gsl/gsl\_matrix.h>

#### **Functions**

double dunn index (gsl matrix \*centroids, int n clusters, gsl matrix \*\*clusters)

#### 5.2.1 Function Documentation

5.2.1.1 double dunn\_index ( gsl\_matrix \* centroids, int n\_clusters, gsl\_matrix \*\* clusters )

Calculates the Dunn Index, a metric for evaluating the clustering results.

#### **Parameters**

centroids	Pointer to matrix containing the centroids
n_clusters	The number of clusters
clusters	Pointer to array of matrices containing data in clusters

#### Returns

The Dunn Index

Definition at line 32 of file fitness.c.

#### 5.3 include/io.h File Reference

```
#include <gsl/gsl_matrix.h>
```

#### **Functions**

- int load\_data (char \*input, gsl\_matrix \*data)
- int save\_results (char \*output, char \*output2, char \*output3, int size, double fitness[size], gsl\_matrix \*\*\*population, int n\_clusters, gsl\_matrix \*\*\*clusters)

#### 5.3.1 Function Documentation

5.3.1.1 int load\_data ( char \* input, gsl\_matrix \* data )

Loads the data from as CSV file into a matrix,

#### **Parameters**

input	Path to the data file
data	Pointer to the GSL matrix to be populated

#### Returns

The status code, 0 for SUCCESS, 1 for ERROR

Definition at line 123 of file io.c.

5.3.1.2 int save\_results ( char \* output, char \* output2, char \* output3, int size, double fitness[size], gsl\_matrix \*\* population, int n\_clusters, gsl\_matrix \*\*\* clusters )

Save the chromosome and fitness value if they are better than previous.

#### **Parameters**

output	Path to save the optimal fitness value
output2	Path to save the optimal fitness centroids
output3	Path to save the optimal cluster results
size	Size of the populations
fitness	Pointer to array of fitness values for the population
population	Population of all chromosomes
n_clusters	The number of clusters
clusters	The clusters for each chromosome in the population

#### Returns

The status code, 0 for SUCCESS, 1 for ERROR

Definition at line 28 of file io.c.

### 5.4 include/operators.h File Reference

```
#include <gsl/gsl_matrix.h>
#include "pcg_basic.h"
```

#### **Functions**

- void crossover (gsl\_matrix \*parent1, gsl\_matrix \*parent2, pcg32\_random\_t \*rng)
- void mutate (gsl\_matrix \*chromosome, gsl\_matrix \*bounds, pcg32\_random\_t \*rng)

#### 5.4.1 Function Documentation

5.4.1.1 void crossover (  $gsl_matrix * parent1$ ,  $gsl_matrix * parent2$ ,  $pcg32_random_t * rng$  )

Performs chromosome crossover by randomly selecting a crossover point and randomly either swapping the top or the bottom half.

#### **Parameters**

len	The length of the chromosome
parent1	The first parent chromosome
parent2	The second parent chromosome
rng	Pointer to the random number generator

Definition at line 34 of file operators.c.

Here is the call graph for this function:



5.4.1.2 void mutate (  $gsl_matrix * chromosome$ ,  $gsl_matrix * bounds$ ,  $pcg32_random_t * rng$  )

Performs mutation, selects a random row and column in the chromosome and mutates it to a random value within the min/max bounds.

#### **Parameters**

chromosome	The chromosome
bounds	The min/max bounds for each dimensions of the data
rng	Pointer to the random number generator

Definition at line 106 of file operators.c.

Here is the call graph for this function:



### 5.5 include/pcg\_basic.h File Reference

#include <inttypes.h>

#### **Data Structures**

• struct pcg\_state\_setseq\_64

#### **Macros**

• #define PCG32\_INITIALIZER { 0x853c49e6748fea9bULL, 0xda3e39cb94b95bdbULL }

#### **Typedefs**

typedef struct pcg\_state\_setseq\_64 pcg32\_random\_t

#### **Functions**

- void pcg32\_srandom (uint64\_t initstate, uint64\_t initseq)
- void pcg32\_srandom\_r (pcg32\_random\_t \*rng, uint64\_t initstate, uint64\_t initseq)
- uint32\_t pcg32\_random (void)
- uint32\_t pcg32\_random\_r (pcg32\_random\_t \*rng)
- uint32\_t pcg32\_boundedrand (uint32\_t bound)
- uint32\_t pcg32\_boundedrand\_r (pcg32\_random\_t \*rng, uint32\_t bound)

#### 5.5.1 Macro Definition Documentation

5.5.1.1 #define PCG32\_INITIALIZER { 0x853c49e6748fea9bULL, 0xda3e39cb94b95bdbULL }

Definition at line 49 of file pcg\_basic.h.

#### 5.5.2 Typedef Documentation

5.5.2.1 typedef struct pcg\_state\_setseq\_64 pcg32\_random\_t

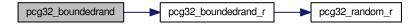
Definition at line 45 of file pcg\_basic.h.

#### 5.5.3 Function Documentation

5.5.3.1 uint32\_t pcg32\_boundedrand ( uint32\_t bound )

Definition at line 112 of file pcg\_basic.c.

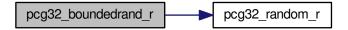
Here is the call graph for this function:



5.5.3.2 uint32\_t pcg32\_boundedrand\_r ( pcg32\_random\_t \* rng, uint32\_t bound )

Definition at line 79 of file pcg\_basic.c.

Here is the call graph for this function:



5.5.3.3 uint32\_t pcg32\_random (void)

Definition at line 69 of file pcg\_basic.c.

Here is the call graph for this function:



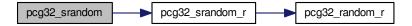
5.5.3.4 uint32\_t pcg32\_random\_r ( pcg32\_random\_t \* rng )

Definition at line 60 of file pcg\_basic.c.

5.5.3.5 void pcg32\_srandom ( uint64\_t initstate, uint64\_t initseq )

Definition at line 51 of file pcg basic.c.

Here is the call graph for this function:



5.5.3.6 void pcg32\_srandom\_r ( pcg32\_random\_t \* rng, uint64\_t initstate, uint64\_t initseq )

Definition at line 42 of file pcg\_basic.c.

Here is the call graph for this function:



#### 5.6 include/selection.h File Reference

#include "pcg\_basic.h"

#### **Functions**

- void gen\_probability (int size, double fitness[size], double probability[size])
- int select\_parent (int size, double probability[size], pcg32\_random\_t \*rng)

#### 5.6.1 Function Documentation

5.6.1.1 void gen\_probability (int size, double fitness[size], double probability[size])

Perform the roulette wheel probability selection, an array is populated with the index of the chromosome to select with a frequency based on the fitness value.

#### **Parameters**

	size	The size of the population
ĺ	fitness	Pointer to an array of fitness values for population
Ì	probability	Pointer to the probability array, populated by function

Definition at line 29 of file selection.c.

5.6.1.2 int select\_parent (int size, double probability[size], pcg32 random t \* rng)

Selects a parent from the population at random with a probability of being selected based on the proabilities provided.

#### **Parameters**

size	The size of the population
probability	Probabilities of each chromosome in population being selected
rng	Pointer to the random number generator

#### Returns

The index of the parent in the population to select

Definition at line 67 of file selection.c.

Here is the call graph for this function:



### 5.7 include/utility.h File Reference

#### Macros

- #define RED "\x1b[31m"
- #define GREEN "\x1b[32m"
- #define YELLOW "\x1b[33m"
- #define BLUE "\x1b[34m"
- #define MAGENTA "\x1b[35m"
- #define CYAN "\x1b[36m"
- #define RESET "\x1b[0m"

#### **Enumerations**

enum debug\_code {
 DEBUG\_CONFIG = 1, DEBUG\_DATA = 2, DEBUG\_CLUSTER = 3, DEBUG\_BOUNDS = 4,
 DEBUG\_CENTROIDS = 5, DEBUG\_DUNN = 6, DEBUG\_CROSSOVER = 7, DEBUG\_MUTATE = 8,
 DEBUG\_PROBABILITY = 9 }

Enumeration of the DEBUG codes.

enum error\_code { SUCCESS = 0, ERROR = 1 }
 Error codes.

#### **Variables**

- int DEBUG
- int VERBOSE

#### 5.7.1 Macro Definition Documentation

5.7.1.1 #define BLUE "\x1b[34m"

Definition at line 27 of file utility.h.

5.7.1.2 #define CYAN "\x1b[36m"

Definition at line 29 of file utility.h.

5.7.1.3 #define GREEN "\x1b[32m"

Definition at line 25 of file utility.h.

5.7.1.4 #define MAGENTA "\x1b[35m"

Definition at line 28 of file utility.h.

5.7.1.5 #define RED "\x1b[31m"

Definition at line 24 of file utility.h.

5.7.1.6 #define RESET "\x1b[0m"

Definition at line 30 of file utility.h.

5.7.1.7 #define YELLOW "\x1b[33m"

Definition at line 26 of file utility.h.

#### 5.7.2 Enumeration Type Documentation

5.7.2.1 enum debug\_code

Enumeration of the DEBUG codes.

#### Enumerator

**DEBUG\_CONFIG** Print all the values parsed from the config file

**DEBUG\_DATA** Print the contents of the data file

**DEBUG\_CLUSTER** Debug the clutering process using lloyd's

**DEBUG\_BOUNDS** Debug the min/max bounds for each dimension

```
    DEBUG_CENTROIDS Debug the randomly generated initial centroids
    DEBUG_DUNN Debug the Dunn Index calculations
    DEBUG_CROSSOVER Debug the crossover operator
    DEBUG_MUTATE Debug the mutation operator
    DEBUG_PROBABILITY Debug output for the probability generation
```

Definition at line 39 of file utility.h.

```
5.7.2.2 enum error_code
```

Error codes.

Enumerator

**SUCCESS** Successful execution **ERROR** Generic error code

Definition at line 56 of file utility.h.

#### 5.7.3 Variable Documentation

5.7.3.1 int DEBUG

Definition at line 40 of file emeans.c.

5.7.3.2 int VERBOSE

Definition at line 40 of file emeans.c.

#### 5.8 README.md File Reference

#### 5.9 src/cluster.c File Reference

```
#include <stdio.h>
#include <stdlib.h>
#include <math.h>
#include <float.h>
#include <stdint.h>
#include <stdint.h>
#include <string.h>
#include <gsl/gsl_matrix.h>
#include <gsl/gsl_blas.h>
#include <gsl/gsl_statistics.h>
#include "utility.h"
#include "pcg_basic.h"
#include "cluster.h"
```

#### **Functions**

- int lloyd\_random (int trials, gsl\_matrix \*data, int n\_clusters, gsl\_matrix \*\*clusters, pcg32\_random\_t \*rng)
- int lloyd\_defined (int trials, gsl\_matrix \*centroids, gsl\_matrix \*data, int n\_clusters, gsl\_matrix \*\*clusters)

- int calc\_centroids (gsl\_matrix \*centroids, gsl\_matrix \*data, int n\_clusters, gsl\_matrix \*\*clusters)
- int calc\_bounds (gsl\_matrix \*data, gsl\_matrix \*bounds)
- int random\_centroids (gsl\_matrix \*centroids, gsl\_matrix \*bounds, pcg32\_random\_t \*rng)

#### 5.9.1 Function Documentation

5.9.1.1 int calc\_bounds ( gsl\_matrix \* data, gsl\_matrix \* bounds )

Calculates the minimum and maximum bounds based on the data.

#### **Parameters**

data	Point to matrix containing the data
bounds	The min/max bounds for each dimensions of the data

#### Returns

The status code, 0 for SUCCESS, 1 for ERROR

Definition at line 334 of file cluster.c.

5.9.1.2 int calc\_centroids ( gsl\_matrix \* centroids, gsl\_matrix \* data, int n\_clusters, gsl\_matrix \*\* clusters )

Calculate the new centroids using the clustering assignment.

#### **Parameters**

centroids	Pointer to matrix containing centroids to be updated
data	Point to matrix containing the data
n_clusters	The number of clusters
cluster	Pointer to vector containing cluster assignment

#### Returns

The status code, 0 for SUCCESS, 1 for ERROR

Definition at line 307 of file cluster.c.

5.9.1.3 int lloyd\_defined ( int trials, gsl\_matrix \* centroids, gsl\_matrix \* data, int n\_clusters, gsl\_matrix \*\* clusters )

Performs Lloyd's algorithm using the defined centroids.

#### **Parameters**

trials	Number of trials to perform
centroids	Pointer to matrix containing the centroids
data	Pointer to matrix containing the data
n_clusters The number of clusters	
clusters	Pointer to array of matrices containing data in clusters

#### Returns

The status code, 0 for SUCCESS, 1 for ERROR

Definition at line 191 of file cluster.c.

Here is the call graph for this function:



5.9.1.4 int lloyd\_random ( int trials, gsl\_matrix \* data, int n\_clusters, gsl\_matrix \*\* clusters, pcg32\_random\_t \* rng )

Performs Lloyd's algorithm using random initial centroids.

#### **Parameters**

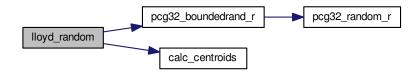
trials	Number of trials to perform	
data Pointer to matrix containing the data		
n_clusters	n_clusters The number of clusters	
clusters	Pointer to array of matrices containing data in clusters	
rng	Pointer to the random number generator	

#### Returns

The status code, 0 for SUCCESS, 1 for ERROR

Definition at line 33 of file cluster.c.

Here is the call graph for this function:



5.9.1.5 int random\_centroids ( gsl\_matrix \* centroids, gsl\_matrix \* bounds, pcg32\_random\_t \* rng )

Generates random centroids within the bounds for each dimension.

#### **Parameters**

centroids	Pointer to matrix containing centroids to be updated	
bounds	The min/max bounds for each dimensions of the data	

rng Pointer to the random number generator

#### Returns

The status code, 0 for SUCCESS, 1 for ERROR

Definition at line 362 of file cluster.c.

Here is the call graph for this function:



#### 5.10 src/emeans.c File Reference

```
#include <stdio.h>
#include <stdlib.h>
#include <stdint.h>
#include <stdbool.h>
#include <string.h>
#include <math.h>
#include <unistd.h>
#include <time.h>
#include <confuse.h>
#include <gsl/gsl_matrix.h>
#include "utility.h"
#include "pcg_basic.h"
#include "io.h"
#include "cluster.h"
#include "fitness.h"
#include "operators.h"
#include "selection.h"
```

#### **Functions**

- int emeans (void)
- int main (int argc, char \*argv[])

#### **Variables**

- int DEBUG
- int VERBOSE
- int64\_t n\_clusters = 3
- int64\_t trials = 1
- int64\_t size = 100
- double m\_rate = 0.01

```
• double c_rate = 0.70
```

- int64\_t max\_iter = 10000
- int64\_t data\_rows = 0
- int64 t data cols = 0
- char \* data file = NULL
- char \* centroids\_file = NULL
- char \* fitness\_file = NULL
- char \* cluster\_file = NULL
- cfg\_opt\_t opts []
- cfg\_t \* cfg

#### 5.10.1 Function Documentation

#### 5.10.1.1 int emeans ( void )

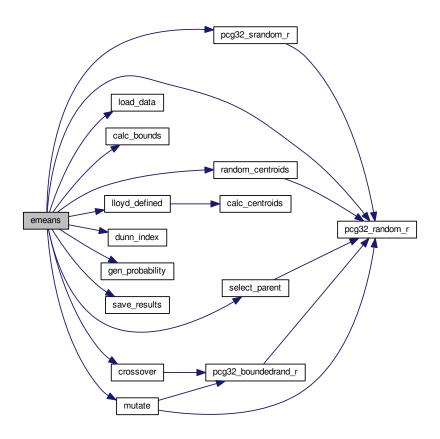
The E-means algorithm, uses a genetic algorithm to optimize the parameters for the K-means implementation of clustering based Lloyds clustering algorithm.

#### Returns

Status code, 0 for SUCCESS, 1 for ERROR

Definition at line 81 of file emeans.c.

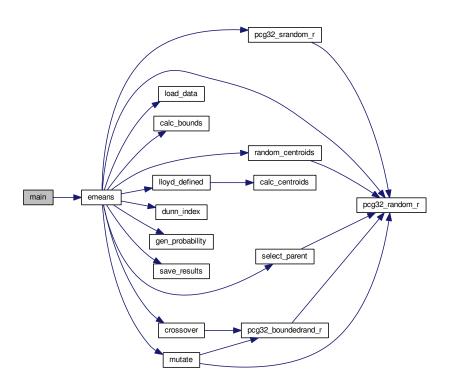
Here is the call graph for this function:



5.10.1.2 int main ( int *argc*, char \* *argv*[] )

Definition at line 241 of file emeans.c.

Here is the call graph for this function:



#### 5.10.2 Variable Documentation

5.10.2.1 double c\_rate = 0.70

Definition at line 47 of file emeans.c.

5.10.2.2 char \* centroids\_file = NULL

Definition at line 52 of file emeans.c.

5.10.2.3 cfg\_t\* cfg

Definition at line 72 of file emeans.c.

5.10.2.4 char \* cluster\_file = NULL

Definition at line 54 of file emeans.c.

5.10.2.5 int64\_t data\_cols = 0

Definition at line 50 of file emeans.c.

```
5.10.2.6 char* data_file = NULL
```

Definition at line 51 of file emeans.c.

```
5.10.2.7 int64_t data_rows = 0
```

Definition at line 49 of file emeans.c.

5.10.2.8 int DEBUG

Definition at line 40 of file emeans.c.

```
5.10.2.9 char * fitness_file = NULL
```

Definition at line 53 of file emeans.c.

5.10.2.10 double m\_rate = 0.01

Definition at line 46 of file emeans.c.

```
5.10.2.11 int64_t max_iter = 10000
```

Definition at line 48 of file emeans.c.

5.10.2.12 int64\_t n\_clusters = 3

Definition at line 43 of file emeans.c.

5.10.2.13 cfg\_opt\_t opts[]

#### Initial value:

```
= {
    CFG_SIMPLE_INT("n_clusters", &n_clusters),
    CFG_SIMPLE_INT("trials", &trials),
    CFG_SIMPLE_INT("size", &size),
    CFG_SIMPLE_FLOAT("m_rate", &m_rate),
    CFG_SIMPLE_FLOAT("c_rate", &c_rate),
    CFG_SIMPLE_INT("max_iter", &max_iter),
    CFG_SIMPLE_INT("data_rows", &data_rows),
    CFG_SIMPLE_INT("data_cols", &data_cols),
    CFG_SIMPLE_STR("data_file", &data_file),
    CFG_SIMPLE_STR("centroids_file", &centroids_file),
    CFG_SIMPLE_STR("fitness_file", &fitness_file),
    CFG_SIMPLE_STR("cluster_file", &cluster_file),
}
```

Definition at line 57 of file emeans.c.

```
5.10.2.14 int64_t size = 100
```

Definition at line 45 of file emeans.c.

```
5.10.2.15 int64_t trials = 1
```

Definition at line 44 of file emeans.c.

5.10.2.16 int VERBOSE

Definition at line 40 of file emeans.c.

#### 5.11 src/fitness.c File Reference

```
#include <stdio.h>
#include <stdlib.h>
#include <math.h>
#include <float.h>
#include <stdint.h>
#include <stdint.h>
#include <string.h>
#include <gsl/gsl_matrix.h>
#include <gsl/gsl_blas.h>
#include <gsl/gsl_statistics.h>
#include "utility.h"
#include "fitness.h"
```

#### **Functions**

• double dunn\_index (gsl\_matrix \*centroids, int n\_clusters, gsl\_matrix \*\*clusters)

#### 5.11.1 Function Documentation

```
5.11.1.1 double dunn_index ( gsl_matrix * centroids, int n_clusters, gsl_matrix ** clusters )
```

Calculates the Dunn Index, a metric for evaluating the clustering results.

#### **Parameters**

centroids Pointer to matrix containing the centroids	
n_clusters The number of clusters	
clusters	Pointer to array of matrices containing data in clusters

#### Returns

The Dunn Index

Definition at line 32 of file fitness.c.

#### 5.12 src/io.c File Reference

```
#include <stdio.h>
#include <stdlib.h>
#include <stdint.h>
#include <float.h>
#include "utility.h"
#include "io.h"
```

#### **Functions**

- int save\_results (char \*output, char \*output2, char \*output3, int size, double fitness[size], gsl\_matrix \*\*population, int n\_clusters, gsl\_matrix \*\*\*clusters)
- int load\_data (char \*input, gsl\_matrix \*data)

#### 5.12.1 Function Documentation

5.12.1.1 int load\_data ( char \* input, gsl\_matrix \* data )

Loads the data from as CSV file into a matrix,

#### **Parameters**

input	Path to the data file
data	Pointer to the GSL matrix to be populated

#### Returns

The status code, 0 for SUCCESS, 1 for ERROR

Definition at line 123 of file io.c.

5.12.1.2 int save\_results ( char \* output, char \* output2, char \* output3, int size, double fitness[size], gsl\_matrix \*\* population, int n\_clusters, gsl\_matrix \*\*\* clusters )

Save the chromosome and fitness value if they are better than previous.

#### **Parameters**

output	Path to save the optimal fitness value
output2	Path to save the optimal fitness centroids
output3	Path to save the optimal cluster results
size	Size of the populations
fitness	Pointer to array of fitness values for the population
population	Population of all chromosomes
n_clusters	The number of clusters
clusters	The clusters for each chromosome in the population

#### Returns

The status code, 0 for SUCCESS, 1 for ERROR

Definition at line 28 of file io.c.

### 5.13 src/operators.c File Reference

```
#include <stdio.h>
#include <stdlib.h>
#include <math.h>
#include <float.h>
#include <stdint.h>
#include <stdint.h>
#include <stdbool.h>
#include <gsl/gsl_matrix.h>
#include <gsl/gsl_blas.h>
#include <gsl/gsl_statistics.h>
#include "utility.h"
#include "fitness.h"
#include "operators.h"
```

#### **Functions**

- void crossover (gsl\_matrix \*parent1, gsl\_matrix \*parent2, pcg32\_random\_t \*rng)
- void mutate (gsl\_matrix \*chromosome, gsl\_matrix \*bounds, pcg32\_random\_t \*rng)

#### 5.13.1 Function Documentation

```
5.13.1.1 void crossover ( gsl_matrix * parent1, gsl_matrix * parent2, pcg32_random_t * rng )
```

Performs chromosome crossover by randomly selecting a crossover point and randomly either swapping the top or the bottom half.

#### **Parameters**

len	The length of the chromosome
parent1	The first parent chromosome
parent2	The second parent chromosome
rng	Pointer to the random number generator

Definition at line 34 of file operators.c.

Here is the call graph for this function:



5.13.1.2 void mutate (  $gsl_matrix * chromosome$ ,  $gsl_matrix * bounds$ ,  $pcg32_random_t * rng$  )

Performs mutation, selects a random row and column in the chromosome and mutates it to a random value within the min/max bounds.

#### **Parameters**

chromosome	chromosome The chromosome	
bounds	bounds The min/max bounds for each dimensions of the data	
rng	Pointer to the random number generator	

Definition at line 106 of file operators.c.

Here is the call graph for this function:



### 5.14 src/pcg\_basic.c File Reference

#include "pcg\_basic.h"

#### **Functions**

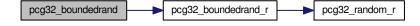
- void pcg32\_srandom\_r (pcg32\_random\_t \*rng, uint64\_t initstate, uint64\_t initseq)
- void pcg32\_srandom (uint64\_t seed, uint64\_t seq)
- uint32\_t pcg32\_random\_r (pcg32\_random\_t \*rng)
- uint32\_t pcg32\_random ()
- uint32\_t pcg32\_boundedrand\_r (pcg32\_random\_t \*rng, uint32\_t bound)
- uint32\_t pcg32\_boundedrand (uint32\_t bound)

#### 5.14.1 Function Documentation

5.14.1.1 uint32\_t pcg32\_boundedrand ( uint32\_t bound )

Definition at line 112 of file pcg\_basic.c.

Here is the call graph for this function:



5.14.1.2 uint32\_t pcg32\_boundedrand\_r ( pcg32\_random\_t \* rng, uint32\_t bound )

Definition at line 79 of file pcg\_basic.c.

Here is the call graph for this function:



5.14.1.3 uint32\_t pcg32\_random ( void )

Definition at line 69 of file pcg\_basic.c.

Here is the call graph for this function:



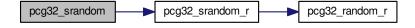
5.14.1.4 uint32\_t pcg32\_random\_r ( pcg32\_random\_t \* rng )

Definition at line 60 of file pcg\_basic.c.

5.14.1.5 void pcg32\_srandom ( uint64\_t seed, uint64\_t seq )

Definition at line 51 of file pcg\_basic.c.

Here is the call graph for this function:



5.14.1.6 void pcg32\_srandom\_r ( pcg32\_random\_t \* rng, uint64\_t initstate, uint64\_t initseq )

Definition at line 42 of file pcg\_basic.c.

Here is the call graph for this function:



#### 5.15 src/selection.c File Reference

```
#include <stdio.h>
#include <stdlib.h>
#include <math.h>
#include <string.h>
#include <stdbool.h>
#include "pcg_basic.h"
#include "utility.h"
#include "selection.h"
```

#### **Functions**

- void gen\_probability (int size, double fitness[size], double probability[size])
- int select\_parent (int size, double probability[size], pcg32\_random\_t \*rng)

#### 5.15.1 Function Documentation

5.15.1.1 void gen\_probability ( int size, double fitness[size], double probability[size] )

Perform the roulette wheel probability selection, an array is populated with the index of the chromosome to select with a frequency based on the fitness value.

#### **Parameters**

size	size The size of the population	
fitness	fitness Pointer to an array of fitness values for population	
probability Pointer to the probability array, populated by function		

Definition at line 29 of file selection.c.

5.15.1.2 int select\_parent ( int size, double probability[size], pcg32\_random\_t \* rng )

Selects a parent from the population at random with a probability of being selected based on the proabilities provided.

#### **Parameters**

size	size The size of the population	
probability Probabilities of each chromosome in population being selected		
rng	Pointer to the random number generator	

#### Returns

The index of the parent in the population to select

Definition at line 67 of file selection.c.

Here is the call graph for this function:



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